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SEQUENCE LISTING

5	<pre><110> Microbiological Research Authority HALLIS, Bassam SILMAN, Nigel SHONE, Clifford Charles SUTTON, John Mark</pre>
10	<120> Delivery of Superoxide Dismutase to Neuronal Cells
	<130> 20994-SOD-heavy chain conjugates
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	<150> GB 9824282.9 <151> 1998-11-05
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40	Leu Gln Asn Lys Ser Leu Glu Glu Leu Leu Ser Asn Leu Glu Ala Leu 50 55 60
	Pro Glu Ser Ile Arg Thr Ala Val Arg Asn Asn Gly Gly His Ala 65 70 75 80
45	Asn His Ser Leu Phe Trp Thr Ile Leu Ser Pro Asn Gly Gly Glu 85 90 95
50	Pro Thr Gly Glu Leu Ala Glu Ala Ile Asn Lys Lys Phe Gly Ser Phe 100 105 110
	Thr Ala Phe Lys Asp Glu Phe Ser Lys Ala Ala Ala Gly Arg Phe Gly 115 120 125
55	Ser Gly Trp Ala Trp Leu Val Val Asn Asn Gly Glu Leu Glu Ile Thr 130 135 140
	Ser Thr Pro Asn Gln Asp Ser Pro Ile Met Glu Gly Lys Thr Pro Ile 145 150 155 160
60	Leu Gly Leu Asp Val Trp Glu His Ala Tyr Tyr Leu Lys Tyr Gln Asn 165 170 175
65	Arg Arg Pro Glu Tyr Ile Ala Ala Phe Trp Asn Ile Val Asn Trp Asp 180 185 190
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15	Asn Thr Tyr Val Thr Asn Leu Asn Ala Ala Leu Glu Gly Hi 35 40 45	is Pro Asp
20	Leu Gln Asn Lys Ser Leu Glu Glu Leu Leu Ser Asn Leu Gl 50 55 60	lu Ala Leu
	Pro Glu Ser Ile Arg Thr Ala Val Arg Asn Asn Gly Gly G 65 70 75	ly His Ala 80
25	Asn His Ser Leu Phe Trp Thr Ile Leu Ser Pro Asn Gly G 85 90	ly Gly Glu 95
	Pro Thr Gly Glu Leu Ala Asp Ala Ile Asn Lys Lys Phe G	ly Ser Phe 10
30	Thr Ala Phe Lys Asp Glu Phe Ser Lys Ala Ala Ala Gly A 115 120 125	rg Phe Gly
	Ser Gly Trp Ala Trp Leu Val Val Asn Asn Gly Glu Leu G 130 135 140	lu Ile Thr
35	Ser Thr Pro Asn Gln Asp Ser Pro Ile Met Glu Gly Lys T 145 150 155	hr Pro Ile 160
40	Leu Gly Leu Asp Val Trp Glu His Ala Tyr Tyr Leu Lys T 165 170	yr Gln Asn 175
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	Leu	Gln 50	Asn	Lys	Ser	Leu	Glu 55	Glu	Leu	Leu	Ser	Asn 60	Leu	Glu	Ala	Leu
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	Asn	His	Ser	Leu	Phe 85	Trp	Thr	Ile	Leu	Ser 90	Pro	Asn	Gly	Gly	Gly 95	Glu
10	Pro	Thr	Gly	Glu 100	Leu	Ala	Asp	Ala	Ile 105	Asn	Lys	Lys	Phe	Gly 110	Ser	Phe
4 -	Thr	Ala	Phe 115	Lys	Asp	Glu	Phe	Ser 120	Lys	Ala	Ala	Ala	Gly 125	Arg	Phe	Gly
15	Ser	Gly 130	Trp	Ala	Trp	Leu	Val 135	Val	Asn	Asn	Gly	Glu 140	Leu	Glu	Ile	Thr
20	Ser 145	Thr	Pro	Asn	Gln	Asp 150	Ser	Pro	Ile	Met	Glu 155	Gly	Lys	Thr	Pro	Ile 160
	Leu	Gly	Leu	Asp	Val 165	Trp	Glu	His	Ala	Tyr 170	Tyr	Leu	Lys	Tyr	Gln 175	Asn
25	Arg	Arg	Pro	Glu 180	Tyr	Ile	Ala	Ala	Phe 185	Trp	Asn	Val	Val	Asn 190	Trp	Asp
30	Glu	Val	Ala 195	Lys	Arg	Tyr	Ser	Glu 200	Ala	Lys	Ala	Lys	Gln 205	Arg	Ser	Cys
30	Gly	Leu 210	Val	Pro	Arg	Gly	Ser 215	Gly	Pro	Gly	Ser	Ala 220	Leu	Asn	Asp	Leu
35	Cys 225	Ile	Lys	Val	Asn	Asn 230	Trp	Asp	Leu	Phe	Phe 235	Ser	Pro	Ser	Glu	Asp 240
	Asn	Phe	Thr	Asn	Asp 245	Leu	Asn	Lys	Gly	Glu 250	Glu	lle	Thr	Ser	Asp 255	Thr
40	Asn	Ile	Glu	Ala 260	Ala	Glu	Glu	Asn /	Ile 265	Ser	Leu	Asp	Leu	Ile 270	Gln	Gln
45	Tyr	Tyr	Leu 275	Thr	Phe	Asn	Phe	Asp 280	Asn	Glu	Pro	Glu	Asn 285	Ile	Ser	Ile
10	Glu	Asn 290	Leu	Ser	Ser	qaA	Ile 295	Ile	Gly	Gln	Leu	Glu 300	Ĺeu	Met	Pro	Asn
50	Ile 305	Glu	Arg	Phe	Pro	Asn 310	Gly	Lys	Lys	Tyr	Glu 315	Leu	Asp	Lys	Tyr	Thr 320
	Met	Phe	His	Tyr	Leu 325	Arg	Ala	Gln	Glu	Phe 330	Glu	His	Gly	Lys	Ser 335	Arg
55	Ile	Ala	Leu	Thr 340	Asn	Ser	Val	Asn	Glu 345	Ala	Leu	Leu	Asn	Pro 350	Ser	Arg
60	Val	Tyr	Thr 355	Phe	Phe	Ser	Ser	Asp 360		Val	Lys	Lys	Val 365	Asn	Lys	Ala
	Thr	Glu 370	Ala	Ala	Met	Phe	Leu 375	Gly	Trp	Val	Glu	Gln 380	Leu	Val	Tyr	Asp
65	Phe 385	Thr	Asp	Glu	Thr	Ser 390	Glu	Val	Ser	Thr	Thr 395	Asp	Lys	Ile	Ala	Asp 400
	Tle	Thr	Tle	Tle	Tle	Dro	ጥ ኣታታ	TTe	Glv	Pro	Δla	Len	Asn	Tle	G1v	Δen

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					405					410		,			415	
5	Met	Leu	Tyr	Lys 420	Asp	Asp	Phe	Val	Gly 425	Ala	Leu	Ile	Phe	Ser 430	Gly	Ala
J	Val	Ile	Leu 435	Leu	Glu	Phe	Ile	Pro 440	Glu	Ile	Ala	Ile	Pro 445	Val	Leu	Gly
10	Thr	Phe 450	Ala	Leu	Val	Ser	Tyr 455	Ile	Ala	Asn	Lys	Val 460	Leu	Thr	Val	Glr
	465		Asp			470			. *		475					480
15			Tyr		485					490					495	
20			Ile	500					505					510		
			Lys 515					520					525			
25		530	Asn				535					540				
20	545		Ser			550					555		_			560
30			Ser		565					570				_	575	
35		•	Leu	580					585					590		_
			Tyr 595					600					605		_	
40		610	Lys				615					620				
45	625		Tyr			630					635					640
75			Asn		645			-		650					655	
50			Leu Val	660					665		•			670		
	·		675					680					685			
55		690	Glu				695					700				
60	705		Ser		_	710					715					720
		_	Tyr		725			_		730			_		735	
65		-	Met	740					745					750	_	
	GIU	тте	Ile 755	Trp	Inr	ьеп	GIN	760	Inr	GIN	GIU	тте	Lys 765	GIN	arg	val

	Val	Phe 770		Tyr	Ser	Gln	Met 775		Asn	Ile	Ser	Asp 780	Tyr	Ile	Asn	Arg
5	Trp 785	Ile	Phe	Val	Thr	Ile 790	Thr	Asn	Asn	Arg	Leu 795	Asn	Asn	Ser	Lys	Ile 800
	Tyr	Ile	Asn	Gly	Arg 805	Leu	Ile	Asp	Gln	Lys 810	Pro	Ile	Ser	Asn	Leu 815	Gly
10	Asn	Ile	His	Ala 820	Ser	Asn	Asn	Ile	Met 825	Phe	Lys	Leu	Asp	Gly 830	Cys	Arg
15	Asp	Thr	His 835	Arg	Tyr	Ile	Trp	Ile 840	Lys	Tyr	Phe	Asn	Leu 845	Phe	Asp	Lys
	Glu	Leu 850	Asn	Glu	Lys	Glu	Ile 855	Lys	Asp	Leu	Tyr	Asp 860	Asn	Gln	Ser	Asn
20	Ser 865	Gly	Ile	Leu	Lys	Asp 870	Phe	Trp	Gly	Asp	Tyr 875	Leu	Gln	Tyr	Asp	Lys 880
	Pro	Tyr	Tyr	Met	Leu 885	Asn	Leu	Tyr	Asp	Pro 890	Asn	Lys	Tyr	Val	Asp 208	Val
25	Asn	Asn	Val	Gly 900	Ile	Arg	Gly	Tyr	Met 905	Tyr	Leu	Lys	Gly	Pro 910	Arg	Gly
30	Ser	Val	Met 915	Thr	Thr	Asn	Ile	Tyr 920	Leu	Asn	Ser	Ser	Leu 925	Tyr	Arg	Gly
		930					935					940			Asn	
35	945					950					955				Asn	960
4.0					965					970					Lys 975	
40				980					985					990	Val	
45	٠		995				. 1	.000				1	.005	•	Lys	
	1	.010				1	.015				1	.020		_	Phe	
50	1025	5			1	.030			•	1	.035			_		.040
	Gln	Ile	Glu	Arg 1	Ser .045	Ser	Arg	Thr		Gly .050	Cys	Ser	Trp		Phe .055	Ile
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- <212> PRT <213> Artificial Sequence
- 65 <220> <223> Description of Artificial Sequence:construct

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	Asn	Thr	Tyr 35	Val	Thr	Asn	Leu	Asn 40			Leu	Glu	Gly 45	30 His		Asp
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15	Pro 65	Glu	Ser	Ile	Arg	Thr 70	Ala	Val	Arg	Asn	Asn 75	Gly	Gly	Gly	His	Ala 80
	Asn	His	Ser	Leu	Phe 85	Trp	Thr	Ile	Leu	Ser 90	Pro	Asn	Gly	Gly	Gly 95	Glu
20	Pro	Thr	Gly	Glu 100	Leu	Ala	Asp	Ala	Ile 105	Asn	Lys	Lys	Phe	Gly 110	Ser	Phe
25	Thr	Ala	Phe 115	Lys	Asp	Glu	Phe	Ser 120	Lys	Ala	Ala	Ala	Gly 125	Arg	Phe	Gly
		130		Ala			135					140				
30	145			Asn		150					155		_			160
25				Asp	165					170					175	
35				Glu 180				-	185					190	:	
40			195	Lys				200					205			- ,
		210		Pro			215					220	•		_	
45	225			Val		230					235			_		240
50				Asp	245		•			250					255	
				Tyr 260 Leu					265				•	270		
55			275	Asp				280					285			
		290		Lys			295					300				
60	305			Thr		310					315				_	320
65				Asp	325					330					335	
				340					345		-1-		-1-	350		- 444

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5	Phe	Ala 370	Gly	Trp	Val	Lys	Gln 375	Ile	Val	Asn	Asp	Phe 380		Ile	Glu	Ala
	Asn 385	Lys	Ser	Asn	Thr	Met 390	Asp	Lys	Ile	Ala	Asp 395	Ile	Ser	Leu	Ile	Val
10	Pro	Туг	Ile	Gly	Leu 405	Ala	Leu	Asn	Val	Gly 410	Asn	Glu	Thr	Aļa	Lys 415	Gly
15	Asn	Phe	Glu	Asn 420	Ala	Phe	Glu	Ile	Ala 425	Gly	Ala	Ser	Ile	Leu 430	Leu	Glu
15	Phe	Ile	Pro 435	Glu	Leu	Leu	Ile	Pro 440	Val	Val	Gly	Ala	Phe 445	Leu	Leu	Glu
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	Leu 465	Thr	Lys	Arg	Asn	Glu 470	Lys	Trp	Ser	Asp	Met 475	туг	Gly	Leu	Ile	Val 480
25	Ala	Gln	Trp	Leu	Ser 485	Thr	Val	Asn	Thr	Gln 490	Phe	Tyr	Thr	Ile	Lys 495	Glu
30	Gly	Met	Tyr	Lys 500	Ala	Leu	Asn	Tyr	Gln 505	Ala	Gln	Ala	Leu	Glu 510	Glu	Ile
	Ile	Lys	Tyr 515	Arg	Tyr	Asn	Ile	Tyr 520	Ser	Glu	Lys	Glu	Lys 525	Ser	Asn	Ile
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	Gln 545	Ala.	Ile	Asp	Asn	Ile 550	Asn.	Asn	Phe		Asn 555	Gly	Cys	Ser	Val	Ser 560
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1 5	Phe	Asp	Asn	Thr 580	Leu	Lys	Lys	Asn	Leu 585	Leu	Asn	Tyr	Ile	Asp 590	Glu	Asn
	Lys	Leu	Tyr 595	Leu	Ile	Gly	Ser	Ala 600	Glu	Tyr	Glu	Lys	Ser 605	Lys	Val	Asn
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	705					710					715					720
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J	Asn	Asn	Ser	Gly 740	Trp	Lys	Ile	Ser	Ile 745	Arg	Gly	Asn	Arg	Ile 750	Ile	Trp
10	Thr	Leu	Ile 755	Asp	Ile	Asn	Gly	Lys 760	Thr	Lys	Ser	Val	Phe 765	Phe ·	Glu	Туз
	Asn	11e 770	Arg	Glu	Asp	Ile.	Ser 775	Glu	Tyr	Ile	Asn	Arg 780	Trp	Phe	Phe	Val
15	785			Asn		790					795				_	800
20				Asn	805					810					815	
				Ile 820					825					830		
25			835	Lys				840					845			
		850		Glu			855			*		860				, -
30	865			Gly		870			÷		875					880
35			_	Asn	885					890	•				895	
				Ile 900					905					910		,,,
10			915	Arg				920					925		_	
. =	ГÀЗ	Ser 930	Asn	Ser	Gln	Ser	Ile 935	Asn.	Asp	Asp	Ile	Val 940	Arg	Lys	Glu	Asp
15	945	•		Leu		950					955		_			960
50		-,		Tyr	965		-			970					975	
	Ile	Ser	Asp	Ser 980	Asp	Glu	Phe	Tyr	Asn 985	Thr	Ile	Gln	Ile	Lys 990	Glu	Тух
55			995	Pro		_	1	1000				1	1005	_	_	
		Ser .010	Thr	Asp.	Glu		Gly 1015	Leu	Ile	Gly		His LO20	Arg	Phe	Tyr	Glu
60	Ser 1025		Ile	Val		Glu 1030	Glu	Tyr	Lys		Tyr L035	Phe	Cys	Ile		Lys 1040
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15	Pro	His	Ile	Asp 20	Lys	Glu	Thr	Met	Asn 25	Ile	His	His	Thr	Lys 30	His	His
20	Asn	Thr	Tyr 35	Val	Thr	Asn	Leu	Asn 40	Ala	Ala	Leu	Glu	Gly 45	His	Pro	Ası
	Leu	Gln 50	Asn	Lys	Ser	Leu	Glu 55	Glu	Leu	Leu	Ser	Asn 60	Leu	Glu	Ala	Let
25	Pro 65	Glu	Ser	Ile	Arg	Thr 70	Ala	Val	Arg	Asn	Asn 75	Gly	Gly	Gly	His	Ala 80
	Asn	His	Ser	Leu	Phe 85	Trp	Thr	Ile	Leu	Ser 90	Pro	Asn	Gly	Gly	Gly 95	Gli
30	Pro	Thr	Gly	Glu 100	Leu	Ala	Asp	Ala	Ile 105	Asn	Lys	Lys	Phe	Gly 110	Ser	Phe
35	Thr	Ala	Phe 115	Lys	Asp	Glu	Phe	Ser 120	Lys	Ala	Ala	Ala	Gly 125	Arg	Phe	Gly
	Ser	Gly 130	Trp	Ala	Trp	Leu	Val 135	Val	Asn	Asn	Gly	Glu 140	Leu	Glu	Ile	Thr
40	Ser 145	Thr	Pro	Asn	Gln	Asp 150	Ser	Pro	Ile	Met	Glu 155	Gly	Lys	Thr	Pro	11e
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45	Arg	Arg	Pro	Glu 180	Tyr	Ile	Ala	Ala	Phe 185	Trp	Asn	Val	Val	Asn 190	Trp	Asp
	Glu	Val	Ala 195	Lys	Arg	Tyr	Ser	Glu 200	Ala	Lys	Ala	Lys	Gln 205	Arg	Ser	Суз
	Gly	Leu 210	Val	Pro	Arg	Gly	Ser 215	Gly	Pro	Gly	Ser	Lys 220	Ala	Pro	Pro	Arg
	Leu 225	Сув	Ile	Arg	Val	Asn 230	Asn	Arg	Glu	Leu	Phe 235	Phe	Val	Ala	Ser	Glu 240
	Ser	Ser	Tyr	Asn	Glu 245	Asn	Asp	Ile	Asn	Thr 250	Pro	Lys	Glu	Ile	Asp 255	Asp
	Thr	Thr	Asn	Leu 260	Asn	Asn	Asn	Tyr	Arg 265	Asn	Asn	Leu	Asp	Glu 270	Val	Ile
	Leu	Asp	Tyr 275	Asn	Ser	Glu	Thr	Ile 280	Pro	Gln	Ile	Ser	Asn 285	Gln	Thr	Lev
	зn	Thr 290	Leu	Val	Gln	Asp	Asp 295	Ser	Tyr	Val	Pro	Arg 300	Tyr	Asp	Ser	Asn

	Gl ₃ 309	y Thi	r Sei	r Glı	ı Ile	310	ı Glı	ı His	s Asr	ı Vai	l Vai	l Asp) Le	u Ası	n Vai	l Phe 320
5	Phe	∋ Туз	r Léu	ı His	325	Glr	Lys	val	l Pro	Gl ₁	ı Gly	/ Glu	ı Thi	r Ası	11e 339	e Ser
	Let	ı Thi	: Ser	Ser 340	: Ile	Asp	Thr	Ala	Leu 345	ı Sei	Glu	ı Glu	ı Ser	Glr 350		l Tyr
10	Thr	Phe	Phe 355	Ser	Ser	Glu	Phe	11e 360	Asn	Thr	: Ile	: Asn	Lys 365		Va]	His
15	Ala	Ala 370	Leu	Phe	Ile	Ser	Trp 375	Ile	Asn	Gln	Val	Ile 380	Arg	Asp	Phe	. Thr
, 0	Thr 385	Glu	Ala	Thr	Gln	Lys 390	Ser	Thr	Phe	Asp	Lys 395	Ile	Ala	Asp	Ile	Ser 400
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	Gln	Lys	Glu	Asn 420	Phe	Lys	Glu	Ala	Phe	Glu	Leu	Leu	Gly	Ala 430		Ile
25	Leu	Leu	Glu 435	Phe	Val	Pro	Glu	Leu 440	Leu	Ile	Pro	Thr	Ile 445	Leu	Val	Phe
30	Thr	Ile 450	Lys	Ser	Phe	Ile	Gly 455	Ser	Ser	Glu	Asn	Lys 460	Asn	Lys	Ile	Ile
00	Lys 465	Ala	Ile	Asn	Asn	Ser 470	Leu	Met	Glu	Arg	Glu 475	Thr	Lys	Trp	Lys	Glu 480
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	Phe	Asn	Lys	Arg 500	Lys	Glu	Gln	Met	Tyr 505	Gln	Ala	Leu	Gln	Asn 510	Gln	Val
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4 5	qaA	Glu 530	Arg	Asn	Arg	Leu	Glu 535	Ser	Glu	Tyr	Asn	Ile 540	Asn	Asn	Ile	Arg
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	Ala	Lys	Val	Ser 580	Lys	Leu	Arg	Glu	Tyr 585	Asp	Glu	Gly	Val	Lys 590	Glu	Tyr
55	Leu	Leu	Asp 595	Tyr	Ile	Ser	Glu	His 600	Arg	Ser	Ile	Leu	Gly 605	Asn	Ser	Val
60	Gln	Glu 610	Leu	Asn	Asp	Leu	Val 615	Thr	Ser	Thr	Leu	Asn 620	Asn	Ser	Ile	Pro
00	Phe 625	Glu	Leu	Ser	Ser	Tyr 630	Thr	Asn	Asp	Lys	Ile 635	Leu	Ile	Leu	Tyr	Phe 640
65	Asn	Lys	Leu-	Tyr	Lys 645	Lys	Ile	Lys		Asn 650	Ser	Ile	Leu	Asp	Met 655	Arg
	Tyr	Glu	Asn	Asn	Lys	Phe	Ile	qaA	Ile	Ser	Gly	Tyr	Gly	Ser	Asn	Ile

- 11 -

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		Gly	1le 690	Tyr	Ser	Ser	Lys	695	Ser	Glu	Val	. Asn	11e 700		Glr	Asn	Asn
10		Asp 705	Ile	Ile	Tyr	Asn	Gly 710	Arg	y Tyr	Gln	Asn	715	Ser	Ile	Ser	Phe	Trp 720
						725					730	•				735	
15	•				740					745					750		Ser
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			770					775					780				Asp
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- 30					820	Asp				825					830		
35				835		Asp	٠		840		•			845			
			850			Leu	*	855			٠		860				_
40	•	363				Ser	870					875					880
45						Tyr 885					890				- 12	895	
45					900	Ser				905					910		
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						Arg 965					970					975	
60					980	Leu		•		985					990		_
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	Leu Pro	Tyr P	ro Tyr	Asp	Ala	Leu 40	Glu	Pro	His	Ile	Asp 45	Lys	Glu	Thr
30	Met Asn 50		is His	Thr	Lys 55	His	His	Asn	Thr	Tyr 60	Val	Thr	Asn	Leu
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40	Val Arg		sn Gly 00	Gly	Gly	His	Ala 105	Asn	His	Ser	Leu	Phe 110	Trp	Thr
	Ile Leu	Ser P	ro Asn	Gly	Gly	Gly 120	Glu	Pro	Thr	Gly	Glu 125	Leu	Ala	Asp
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30	Val Asn	Asn G	ly Glu 165	Leu	Glu	Ile	Thr	Ser 170	Thr	Pro	Asn	Gln	Asp 175	Ser
55	Pro Ile		lu Gly 80	Lys	Thr	Pro	Ile 185	Leu	Gly	Leu	Asp	Val 190	Trp	Glu
	His Ala	Tyr T 195	yr Leu	Lys	Tyr	Gln 200	Asn	Arg	Arg	Pro	Glu 205	Tyr	Ile	Ala
60	Ala Phe	-	sn Val	Val	Asn 215	Trp	Asp	Glu	Val	Ala 220	Lys	Arg	Tyr	Ser
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10	Asp 305	Asn	Glu	Pro	Glu	Asn 310	Ile	Ser	Ile	Glu	Asn 315		Ser	Ser	-	Ile 320	
15	Ile	Gly	Gln	Leu	Glu 325	Leu	Met	Pro	Asn	Ile 330	Glu	Arg	Phe	Pro	Asn 335	_	
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,	Val	Ser	Thr	Thr 420	qaA	Lys	Ile	Ala	Asp 425	Ile	Thr	Ile	Ile	Ile 430	Pro	Tyr	
35	Ile	Gly	Pro 435	Ala	Leu	Asn	Ile	Gly 440	Asn	Met	Leu	Tyr	Lys 445	Asp	Asp	Phe	
•	Val	Gly 450	Ala	Leu	Ile	Phe	Ser 455	Gly	Ala	Val	Ile	Leu 460	Leu	Glu	Phe	Ile	
40	Pro 465	Glu	Ile	Ala	Ile	Pro 470	Val	Leu	Gly	Thr	Phe 475	Ala	Leu	Val	Ser	Tyr 480	
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	Ala	Ser	Leu	Lys	Asp	Ala	Leu	Leu	Lys	Tyr	Ile	Tyr	Asp	Asn	Arg	Gly	

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610 615 620 Thr Leu Ile Gly Gln Val Asp Arg Leu Lys Asp Lys Val Asn Asn Thr 630 635 5 Leu Ser Thr Asp Ile Pro Phe Gln Leu Ser Lys Tyr Val Asp Asn Gln 650 Arg Leu Leu Ser Thr Phe Thr Glu Tyr Ile Lys Asn Ile Ile Asn Thr 10 Ser Ile Leu Asn Leu Arg Tyr Glu Ser Asn His Leu Ile Asp Leu Ser 15 Arg Tyr Ala Ser Lys Ile Asn Ile Gly Ser Lys Val Asn Phe Asp Pro Ile Asp Lys Asn Gln Ile Gln Leu Phe Asn Leu Glu Ser Ser Lys Ile 20 Glu Val Ile Leu Lys Asn Ala Ile Val Tyr Asn Ser Met Tyr Glu Asn Phe Ser Thr Ser Phe Trp Ile Arg Ile Pro Lys Tyr Phe Asn Ser Ile 25 Ser Leu Asn Asn Glu Tyr Thr Ile Ile Asn Cys Met Glu Asn Asn Ser 760 30 Gly Trp Lys Val Ser Leu Asn Tyr Gly Glu Ile Ile Trp Thr Leu Gln Asp Thr Gln Glu Ile Lys Gln Arg Val Val Phe Lys Tyr Ser Gln Met 35 Ile Asn Ile Ser Asp Tyr Ile Asn Arg Trp Ile Phe Val Thr Ile Thr 805 810 Asn Asn Arg Leu Asn Asn Ser Lys Ile Tyr Ile Asn Gly Arg Leu Ile 40 820 Asp Gln Lys Pro Ile Ser Asn Leu Gly Asn Ile His Ala Ser Asn Asn 840 45 Ile Met Phe Lys Leu Asp Gly Cys Arg Asp Thr His Arg Tyr Ile Trp Ile Lys Tyr Phe Asn Leu Phe Asp Lys Glu Leu Asn Glu Lys Glu Ile 870 875 50 Lys Asp Leu Tyr Asp Asn Gln Ser Asn Ser Gly Ile Leu Lys Asp Phe 890 Trp Gly Asp Tyr Leu Gln Tyr Asp Lys Pro Tyr Tyr Met Leu Asn Leu 55 Tyr Asp Pro Asn Lys Tyr Val Asp Val Asn Asn Val Gly Ile Arg Gly 920 60 Tyr Met Tyr Leu Lys Gly Pro Arg Gly Ser Val Met Thr Thr Asn Ile Tyr Leu Asn Ser Ser Leu Tyr Arg Gly Thr Lys Phe Ile Ile Lys Lys 65 Tyr Ala Ser Gly Asn Lys Asp Asn Ile Val Arg Asn Asn Asp Arg Val 970 965

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	Туз	Ile	Asn	Val 980	Val	Val	Lys	Asn	Lys 985		Тух	Arg	, Leu	Ala 990		Asn
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	Asp	Val 1010	Gly	Asn	Leu	Ser	Gln 1015	Val	Val	Val		Lys 1020		Lys	Asn	Asp
10	Gln 102	Gly	Ile	Thr	Asn	Lys 1030	Cys	Lys	Met		Leu 1035	Gln	Asp	Asn	Asn	Gly 1040
15	Asn	Asp	Ile	Gly	Phe 1045	Ile	Gly	Phe		Gln 1050	Phe	Asn	Asn	Ile	Ala 1055	-
	Ŀeu	Val	Ala	Ser 1060	Asn	Trp	Tyr	Asn	Arg 1065	Gln	Ile	Glu		Ser 1070		Arg
20	Thr	Leu	Gly 1075	Сув	Ser	Trp	Glu :	Phe 1080	Ile	Pro	Val		Asp 1085	Gly	Trp	Gly
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40		•		20			Arg Ala		25	His	*			30	Pro	
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5	His	Ala	Tyr 195	Tyr	Leu	Lys	Tyr	Gln 200	Asn	Arg	Arg	Pro	Glu 205	Tyr	Ile	Ala
	Ala	Phe 210	Trp	Asn	Val	Val	Asn 215	Trp	qaA	Glu	Val	Ala 220	Lys	Arg	Tyr	Ser
10	Glu 225	Ala	Lys	Ala	Ľys	Gln 230	Arg	Ser	Сув	Gly	Leu 235	Val	Pro	Aṛg	Gly	Ser 240
15	Gly	Pro	Gly	Ser	Lys 245	Ala	Pro	Gly	Ile	Cys 250	Ile	Asp	Val	Asp	Asn 255	Glu
15	Asp	Leu	Phe	Phe 260	Ile	Ala	Asp	Lys	Asn 265	Ser	Phe	Ser	Asp	Asp 270	Leu	Ser
20	Lys	Asn	Glu 275	Arg	Ile	Glu	Tyr	Asn 280	Thr	Gln	Ser	Asn	Tyr 285	Ile	Glu	Asn
	Asp	Phe 290	Pro	Ile	Asn	Glu	Leu 295	Ile	Leu	Asp	Thr	Asp 300	Leu	Ile	Ser	Lys
25	Ile 305	Glu	Leu	Pro	Ser	Glu 310	Asn	Thr	Glu	Ser	Leu 315	Thr	Asp	Phe	Asn	Val 320
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30	Asp	Glu	Asn	Thr 340	Ile	Phe	Gln	Tyr	Leu 345	Tyr	Ser	Gln	Thr	Phe 350	Pro	Leu
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65	Asn	Thr	Gln 515	Phe	Tyr	Thr	Ile	Lys 520	Glu	Gly	Met	Tyr	Lys 525	Ala	Leu	Asn
	Tyr	Gln	Ala	Gln	Ala	Leu	Glu	Glu	Ile	Ile	Lys	Tyr	Arg	Tyr	Asn	Ile

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530 535 540 Tyr Ser Glu Lys Glu Lys Ser Asn Ile Asn Ile Asp Phe Asn Asp Ile 5 Asn Ser Lys Leu Asn Glu Gly Ile Asn Gln Ala Ile Asp Asn Ile Asn Asn Phe Ile Asn Gly Cys Ser Val Ser Tyr Leu Met Lys Lys Met Ile 10 585 Pro Leu Ala Val Glu Lys Leu Leu Asp Phe Asp Asn Thr Leu Lys Lys 15 Asn Leu Leu Asn Tyr Ile Asp Glu Asn Lys Leu Tyr Leu Ile Gly Ser Ala Glu Tyr Gļu Lys Ser Lys Val Asn Lys Tyr Leu Lys Thr Ile Met 630 20 Pro Phe Asp Leu Ser Ile Tyr Thr Asn Asp Thr Ile Leu Ile Glu Met Phe Asn Lys Tyr Asn Ser Glu Ile Leu Asn Asn Ile Ile Leu Asn Leu 25 665 Arg Tyr Lys Asp Asn Asn Leu Ile Asp Leu Ser Gly Tyr Gly Ala Lys 685 30 Val Glu Val Tyr Asp Gly Val Glu Leu Asn Asp Lys Asn Gln Phe Lys Leu Thr Ser Ser Ala Asn Ser Lys Ile Arg Val Thr Gln Asn Gln Asn 35 Ile Ile Phe Asn Ser Val Phe Leu Asp Phe Ser Val Ser Phe Trp Ile 725 735 Arg Ile Pro Lys Tyr Lys Asn Asp Gly Ile Gln Asn Tyr Ile His Asn 40 Glu Tyr Thr Ile Ile Asn Cys Met Lys Asn Asn Ser Gly Trp Lys Ile 45 Ser Ile Arg Gly Asn Arg Ile Ile Trp Thr Leu Ile Asp Ile Asn Gly Lys Thr Lys Ser Val Phe Phe Glu Tyr Asn Ile Arg Glu Asp Ile Ser 790 795 50 Glu Tyr Ile Asn Arg Trp Phe Phe Val Thr Ile Thr Asn Asn Leu Asn Asn Ala Lys Ile Tyr Ile Asn Gly Lys Leu Glu Ser Asn Thr Asp Ile 55 Lys Asp Ile Arg Glu Val Ile Ala Asn Gly Glu Ile Ile Phe Lys Leu 840 60 Asp Gly Asp Ile Asp Arg Thr Gln Phe Ile Trp Met Lys Tyr Phe Ser 855 Ile Phe Asn Thr Glu Leu Ser Gln Ser Asn Ile Glu Glu Arg Tyr Lys 870 65 Ile Gln Ser Tyr Ser Glu Tyr Leu Lys Asp Phe Trp Gly Asn Pro Leu 890

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·	Ser	Lys 930	Tyr	Asn	Gln	Asn	Ser 935	Lys	Tyr	Ile	Asn	Tyr 940	Arg	Asp	Leu	Tyr
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	Asn	Leu	Asn	Gln 980	Glu	Trp	Arg	Val	Tyr 985	Thr	Tyr	Lys	Tyr	Phe 990	Lys	Lys
20	Glu	Glu	Glu 995	Lys	Leu	Phe		Ala LOOO	Pro	Ile	Ser		Ser 1005	Asp	Glu	Phe
		Asn 1010	Thr	Ile	Gln		Lys 1015	Glu	Tyr	qaA		Gln L020	Pro	Thr	Tyr	Ser
25	Cys 1025	Gln 5	Leu	Leu		Lys 1030	Lys	Asp	Glu		Ser 1035	Thr	Asp	Glu		Gly 1040
30	Leu	Ile	Gly		His LO45	Arg	Phe	Tyr		Ser 1050	Gly	Ile	Val		Glu 1055	Glu
	Tyr	Lys		Tyr 1060	Phe	Cys	Ile		Lys 1065	Trp	Tyr	Leu		Glu 1070	Val	Lys
35	Arg	Lys	Pro L075	Tyr	Asn	Leu		Leu .080	Gly	Cys	Asn		Gln 1085	Phe	Ile	Pro
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	Leu	Pro	Tyr 35	Pro	Tyr	Asp	Ala	Leu 40	Glu	Pro	His	Ile	Asp 45	Lys	Glu	Thr
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00	Glu	Leu	Leu	Ser	Asn 85	Leu	Glu	Ala	Leu	Pro 90	Glu	Ser	Ile	Arg	Thr 95	Ala

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•	Ala	Ile 130	Asn	Lys	Lys	Phe	Gly 135	Ser	Phe	Thr	Ala	Phe 140	Lys	Asp	Glu	Phe
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15	Val	Asn	Asn	Gly	Glu 165	Leu	Glu	Ile	Thr	Ser 170	Thr	Pro	Asn	Gln	Asp 175	Ser
13	Pro	Ile	Met	Glu 180	Gly	Lys	Thr	Pro	Ile 185	Leu	Gly	Leu	Asp	Val 190	Trp	Glu
20	His	Ala	Tyr 195	Tyr	Leu	Lys	Tyr	Gln 200	Asn	Arg	Arg	Pro	Glu 205	Tyr	Ile	Ala
	Ala	Phe 210	Trp	Asn	Val	Val	Asn 215	Trp	Asp	Glu	Val	Ala 220	Lys	Arg	Tyr	Ser
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35	Ile	Asn	Thr 275	Pro	Lys	Glu	Ile	Asp 280	Asp	Thr	Thr	Asn	Leu 285	Asn	Asn	Asn
	Tyr	Arg 290	Asn	Asn	Leu	Asp	Glu 295	Val	Ile	Leu	qaA	Tyr 300	Asn	Ser	Glu	Thr
40	Ile 305	Pro	Gln	Ile	Ser	Asn 310	Gln	Thr	Leu	Asn	Thr 315	Leu	Val	Gln	Asp	Asp 320
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50	Val	Pro	Glu 355	Gly	Glu	Thr	Asn	Ile 360	Ser	Leu	Thr	Ser	Ser 365	Ile	Asp	Thr
	Ala	Leu 370	Ser	Glu	Glu	Ser	Gln 375	Val	Tyr	Thr	Phe	Phe 380	Ser	Ser	Glu	Phe
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,	Thr	Phe	Asp	Lys 420	Ile	Ala	Asp	Ile	Ser 425	Leu	Val	Val	Pro	Tyr 430	Val	Gly
65	Leu	Ala	Leu 435	Asn	Ile	Gly	Asn	Glu 440	Val	Gln	Lys	Glu	Asn 445	Phe	Lys	Glu
	Ala	Phe	Glu	Leu	Leu	Gly	Ala	Gly	Ile	Leu	Leu	Glu	Phe	Val	Pro	Glu

		450)				455	5				460)			
5	Leu 465	. Leu	Ile	Pro	Thr	1le 470	Leu	ı Val	. Phe	. Thr	11e 475	Lys	Ser	Phe	: Ile	Gly 480
	Ser	Ser	Glu	Asn	Lys 485	Asn	Lys	Ile	: Ile	Lys 490	Ala	Ile	Asn	Asn	Ser 495	Leu
10				500					505	•				51,0		Ser
4 =			515					520					525	_		Gln
15		530					535					540				Ile
20	545					550			•		555					Glu 560
				Asn	565					570	•			_	575	
25		•		Met 580					585				٠.	590		
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30		910	,	Glu			615				*	620	•			
35	625			Ile		630				•	635				. •	640
				Leu	645	-				650					655	
40	•			660		·			665					670	•	Ile
45			675	Ser				680					685			
45		690		Gly			695					700				
50	705					710					715					Pro 720
				Asn	725					730					735	
55	Tyr			740					745					750		
	Asn		/55					760					765			
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65	Trp 785					790					795					800
	Tyr	Thr	Gln	Met	Ile 805	Ser	Ile	Ser	Asp	Tyr 810	Ile	Asn	Lys	Trp	Ile 815	Phe

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	vaı	1111	116	820	non	non		БСи	825	non	DCI	n. g	116	830	116	ASII
5	Gly	Asn	Leu 835	Ile	Asp	Glu	Lys	Ser 840	Ile	Ser	Asn	Leu	Gly 845	Asp	Ile	His
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15	Thr	Glu	Ile	Glu	Thr 885	Leu	Tyr	Ser	Asp	Glu 890	Pro	Asp	Pro	Ser	Ile 895	Leu
	Lys	Asp	Phe	Trp 900	Gly	Asn	Tyr	Leu	Leu 905	Tyr	Asn	Lys	Arg	Tyr 910	Tyr	Leu
20	Leu	Asn	Leu 915	Leu	Arg	Thr	qaA	Lys 920	Ser	Ile	Thr	Gln	Asn 925	Ser	Asn	Phe
	Leu	Asn 930	Ile	Asn	Gln	Gln	Arg 935	Gly	Val	Tyr	Gl'n	Lys 940	Pro	Asn	Ile	Phe
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ĪĪ	Leu	Ala	Tyr	Ile 980	Asn	Val	Val	Asp	Arg 985		Val	Glu	Tyr	Arg 990	Leu	Tyr
35	Ala	Asp	Ile 995	Ser	Ile	Ala		Pro L000	Glu	Lys	Ile		Lys 1005	Leu	Ile	Arg
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	Tyr	Tyr		Asn 1060	Ile	Arg	Lys		Thr L065	Ser	Ser	Asn		Cys .070	Phe	Trp
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	Leu	Gly	Tyr	Leu 20	Gly	Ser	Arg	Gln	Lys 25	His	Ser	Arg	Gly	Ser 30	Pro	Ala
5	Leu	Pro	Tyr 35	Pro	Tyr	Asp	Ala	Leu 40	Glu	Pro	His	Ile	Asp 45	Lys	Glu	Thr
	Met	Asn 50	Ile	His	His	Thr	Lys 55	His	His	Asn	Thr	Tyr 60	Val	Thr	Asn	Leu
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35	His	Ala	Tyr 195	Tyr	Leu	Lys	Tyr	Gln 200	Asn	Arg	Arg	Pro	Glu 205	Tyr	Ile	Ala
	Ala	Phe 210	Trp	Asn	Val	Val	Asn 215	Trp	Asp	Glu	Val	Ala 220	Гуs	Arg	Tyr	Ser
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		-											•			
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	Description of mitochondrial			modified	humar
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Leu Gly Tyr Leu Gly Ser Arg Gln 20